

1

#10

SEQUENCE LISTING

<110>	MARCHION	NI, MARK	(
	JARPE, M	ICHAEL	
	EBENDAL,	TED	

- <120> METHODS FOR TREATING NEUROLOGICAL INJURIES AND DISORDERS
- <130> 47506(47843)
- <140> 09/756,481
- <141> 2001-01-08
- <150> PCT/US99/15106
- <151> 1999-07-02
- <150> 60/091,791
- <151> 1998-07-06
- <160> 4
- <170> PatentIn Ver. 2.1
- <210> 1
- <211> 1387
- <212> DNA
- <213> Mus sp.
- <220>
- <221> CDS
- <222> (218) .. (1288)
- <400> 1
- cccttctcca gggactctgg ctgccagcag ctccgccttt cagatcaatt ctcgaccacc 60
- caccttggga ctgccgccca gtcctgccct ctggatcagt ggggtccaga cacgccccct 120
- ccaggacete aaagcacece egacetaagg teaccagece actggeecea gacgcagtgg 180
- gctccgctga ctctcttgga cacctcctgg gaggaaa atg ctc cct gtc tgc cat 235

 Met Leu Pro Val Cys His

 1
- cgt ttt tgc gac cac ctc ctc ctc ctg ctc ttg ctg ccc tcg acg acc 283
 Arg Phe Cys Asp His Leu Leu Leu Leu Leu Leu Leu Pro Ser Thr Thr
 10 15 20
- ctg gcc ccc gcg cca gca tcc atg ggc ccc gct gcc gcc ctg ctc cag 331 Leu Ala Pro Ala Pro Ala Ser Met Gly Pro Ala Ala Ala Leu Leu Gln 25 30 35
- gtt ctt ggg ctt ccc gaa gcg ccc cgg agc gtc ccc aca cac cga cct 379 Val Leu Gly Leu Pro Glu Ala Pro Arg Ser Val Pro Thr His Arg Pro 40 45 50

gtg Val 55	cct Pro	cct Pro	gtc Val	atg Met	tgg Trp 60	cgc Arg	cta Leu	ttc Phe	cgt Arg	cgc Arg 65	cgt Arg	gac Asp	ccc Pro	cag Gln	gag Glu 70	427
gcc Ala	aga Arg	gtg Val	gga Gly	cgc Arg 75	cct Pro	ctg Leu	cgg Arg	cca Pro	tgc Cys 80	cac His	gtg Val	gag Glu	gaa Glu	cta Leu 85	gl ^y aaa	475
gtc Val	gcc Ala	gga Gly	aac Asn 90	att Ile	gtg Val	cgc Arg	cac His	atc Ile 95	ccc Pro	gac Asp	agc Ser	ggt Gly	ctg Leu 100	tcc Ser	tcc Ser	523
agg Arg	ccc Pro	gca Ala 105	caa Gln	ccc Pro	gcc Ala	agg Arg	acc Thr 110	tcg Ser	ggg Gly	ctg Leu	tgc Cys	ccc Pro 115	gag Glu	tgg Trp	aca Thr	571
gtc Val	gtc Val 120	ttt Phe	gac Asp	ctg Leu	tcg Ser	aat Asn 125	gtg Val	gag Glu	ccc Pro	aca Thr	gag Glu 130	cgc Arg	cca Pro	aca Thr	cgc Arg	619
gcg Ala 135	cgc Arg	tta Leu	gag Glu	ttg Leu	cgg Arg 140	ctg Leu	gag Glu	gct Ala	gag Glu	tgt Cys 145	gaa Glu	gat Asp	aca Thr	gga Gly	999 Gly 150	667
tgg Trp	gag Glu	cta Leu	agc Ser	gtg Val 155	gca Ala	ctg Leu	tgg Trp	gcc Ala	gac Asp 160	gca Ala	gag Glu	cat His	cca Pro	999 Gly 165	cct Pro	715
gag Glu	ctg Leu	ctg Leu	cgc Arg 170	gtg Val	ccg Pro	gcg Ala	cca Pro	cca Pro 175	gly aaa	gtg Val	ctc Leu	ctg Leu	cgc Arg 180	gca Ala	gac Asp	763
cta Leu	ctg Leu	999 Gly 185	act Thr	gca Ala	gta Val	gcc Ala	gcc Ala 190	aac Asn	gca Ala	tca Ser	gtg Val	ccc Pro 195	tgt Cys	act Thr	gtg Val	811
cgc Arg	ctg Leu 200	gcg Ala	ctg Leu	tca Ser	ctg Leu	cac His 205	cct Pro	gly aaa	gcc Ala	act Thr	gca Ala 210	gcc Ala	tgt Cys	gjà aaa	cgc Arg	859
ctg Leu 215	gct Ala	gag Glu	gcc Ala	tcc Ser	ctg Leu 220	ctg Leu	ctg Leu	gtg Val	acg Thr	ctg Leu 225	gac Asp	cca Pro	cgc Arg	ctg Leu	tgt Cys 230	907
ccc Pro	ttg Leu	ccg Pro	cga Arg	ttg Leu 235	cgg Arg	cgc Arg	cac His	acg Thr	gag Glu 240	ccc Pro	agg Arg	gta Val	gaa Glu	gtt Val 245	ggt Gly	955
cca Pro	gtg Val	ggc Gly	act Thr 250	tgt Cys	cgt Arg	acc Thr	cga Arg	cgg Arg 255	ttg Leu	cat His	gtg Val	agc Ser	ttc Phe 260	cgt Arg	gag Glu	1003
gtg Val	ggc Gly	tgg Trp 265	cac His	cgt Arg	tgg Trp	gtg Val	atc Ile 270	gcg Ala	ccg Pro	cgt Arg	ggc Gly	ttc Phe 275	cta Leu	gcc Ala	aac Asn	1051

Phe	tgc Cys 280	cag Gln	ggc Gly	acg Thr	tgc Cys	gca Ala 285	cta Leu	ccc Pro	gaa Glu	acg Thr	ctg Leu 290	agg Arg	gga Gly	ccc Pro	ggc Gly
999 Gly 295	ccg Pro	cct Pro	gca Ala	ctc Leu	aac Asn 300	cac His	gct Ala	gtg Val	ctg Leu	cgc Arg 305	gcg Ala	ctc Leu	atg Met	cac His	gca Ala 310
gct Ala	gct Ala	ccc Pro	acc Thr	ccg Pro 315	ggt Gly	gca Ala	ggc Gly	tcg Ser	ccc Pro 320	tgc Cys	tgc Cys	gtg Val	cca Pro	gag Glu 325	cgt Arg
				tcc Ser											
				gaa Glu											
tgad	ccaco	ccg g	ggaca	accct	t to	caggo	gacco	g cc	ccaco	gcaa	aago	caggg	gac t	gttt	tgttca
tgti	ttat	tg g	gtgad	caaaa	aa go	cttaa	aaaca	a aat	ttga	act					
<213	0> 2 l> 35 2> PF B> Mu	RТ	o .												
		_													
<400	0> 2			Cvc	Wie	λνα	Dho	Cara	ħ a n	11.1 a	Tou	T 011	Τ α	T	T
<400	0> 2			Cys 5	His	Arg	Phe	Cys	Asp 10	His	Leu	Leu	Leu	Leu 15	Leu
<400 Met 1)> 2 Leu	Pro	Val						10					15	
<400 Met 1 Leu)> 2 Leu Leu	Pro Pro	Val Ser 20	5	Thr	Leu	Ala	Pro 25	10 Ala	Pro	Ala	Ser	Met 30	15 Gly	Pro
<400 Met 1 Leu)> 2 Leu Leu	Pro Pro Ala 35	Val Ser 20 Leu	5 Thr	Thr Gln Pro	Leu Val Val	Ala Leu 40 Pro	Pro 25 Gly	10 Ala Leu Val	Pro Pro	Ala Glu Trp	Ser Ala 45 Arg	Met 30 Pro	15 Gly Arg	Pro Ser
<400 Met 1 Leu Ala Val	Leu Ala Pro 50	Pro Pro Ala 35	Val Ser 20 Leu His	5 Thr Leu Arg	Thr Gln Pro	Leu Val Val 55	Ala Leu 40 Pro	Pro 25 Gly Pro	10 Ala Leu Val	Pro Pro Met	Ala Glu Trp 60	Ser Ala 45 Arg	Met 30 Pro Leu	15 Gly Arg Phe	Pro Ser Arg
<400 Met 1 Leu Ala Val Arg 65	D> 2 Leu Leu Ala Pro 50	Pro Pro Ala 35 Thr	Val Ser 20 Leu His	5 Thr Leu Arg	Thr Gln Pro Glu 70	Leu Val Val 55 Ala	Ala Leu 40 Pro	Pro 25 Gly Pro Val	10 Ala Leu Val Gly	Pro Pro Met Arg	Ala Glu Trp 60 Pro	Ser Ala 45 Arg Leu	Met 30 Pro Leu Arg	15 Gly Arg Phe	Pro Ser Arg Cys 80
<400 Met 1 Leu Ala Val Arg 65 His	D> 2 Leu Leu Ala Pro 50 Arg	Pro Pro Ala 35 Thr Asp	Val Ser 20 Leu His Pro	5 Thr Leu Arg Gln Leu	Thr Gln Pro Glu 70 Gly	Leu Val Val 55 Ala Val	Ala Leu 40 Pro Arg	Pro 25 Gly Pro Val	10 Ala Leu Val Gly Asn 90	Pro Pro Met Arg 75	Ala Glu Trp 60 Pro	Ser Ala 45 Arg Leu Arg	Met 30 Pro Leu Arg	15 Gly Arg Phe Pro Ile 95	Pro Ser Arg Cys 80 Pro
<400 Met 1 Leu Ala Val Arg 65 His	D> 2 Leu Leu Ala Pro 50 Arg Val	Pro Pro Ala 35 Thr Asp Glu Gly	Val Ser 20 Leu His Pro Glu Leu 100	5 Thr Leu Arg Gln Leu 85	Thr Gln Pro Glu 70 Gly Ser	Leu Val Val 55 Ala Val	Ala Leu 40 Pro Arg Ala Pro	Pro 25 Gly Pro Val Gly Ala 105	10 Ala Leu Val Gly Asn 90 Gln	Pro Pro Met Arg 75 Ile	Ala Glu Trp 60 Pro Val	Ser Ala 45 Arg Leu Arg	Met 30 Pro Leu Arg His	15 Gly Arg Phe Pro Ile 95 Ser	Pro Ser Arg Cys 80 Pro

Cys Glu Asp Thr Gly Gly Trp Glu Leu Ser Val Ala Leu Trp Ala Asp 145 150 155 160

Ala Glu His Pro Gly Pro Glu Leu Leu Arg Val Pro Ala Pro Pro Gly
165 170 175

Val Leu Leu Arg Ala Asp Leu Leu Gly Thr Ala Val Ala Ala Asn Ala 180 185 190

Ser Val Pro Cys Thr Val Arg Leu Ala Leu Ser Leu His Pro Gly Ala 195 200 205

Thr Ala Ala Cys Gly Arg Leu Ala Glu Ala Ser Leu Leu Leu Val Thr 210 215 220

Leu Asp Pro Arg Leu Cys Pro Leu Pro Arg Leu Arg Arg His Thr Glu 225 230 235 240

Pro Arg Val Glu Val Gly Pro Val Gly Thr Cys Arg Thr Arg Arg Leu 245 250 255

His Val Ser Phe Arg Glu Val Gly Trp His Arg Trp Val Ile Ala Pro 260 265 270

Arg Gly Phe Leu Ala Asn Phe Cys Gln Gly Thr Cys Ala Leu Pro Glu 275 280 285

Thr Leu Arg Gly Pro Gly Gly Pro Pro Ala Leu Asn His Ala Val Leu 290 295 300

Arg Ala Leu Met His Ala Ala Ala Pro Thr Pro Gly Ala Gly Ser Pro 305 310 315 320

Cys Cys Val Pro Glu Arg Leu Ser Pro Ile Ser Val Leu Phe Phe Asp 325 330 335

Asn Ser Asp Asn Val Val Leu Arg His Tyr Glu Asp Met Val Val Asp 340 345 350

Glu Cys Gly Cys Arg 355

<210> 3

<211> 27

<212> DNA

<213> Mus sp.

<400> 3

gcagccacac tectecacca ceatgtt

<210> 4

<211> 9

<212> PRT

<213> Mus sp.

27

<400> 4
Asn Met Val Val Glu Glu Cys Gly Cys
1 5